90

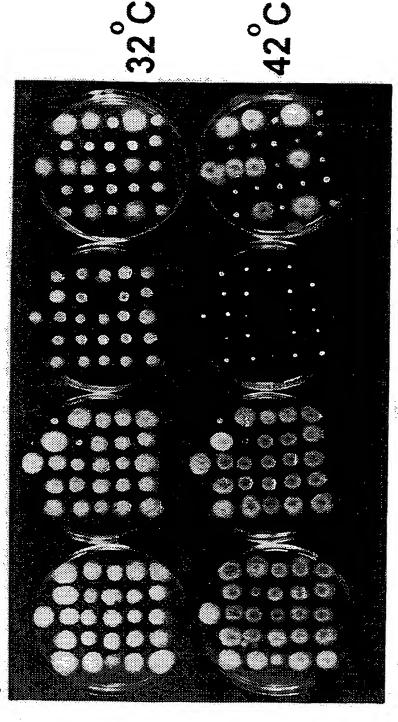
HUM MGGEQEEERFDGMLLAMAQQHEGGVQELVNTFFSFLRRKTDFFIGGEEGMAEKLITQTFSHHNQLAQKTRREKRARQEAERREKAERAAR

rat mggegeerfdgmllamaqqheggvgelvntffsflrrtdffiggeegmaeklitqtfmhhqlaqk**a**rrekrarqe*t*errekaeraar

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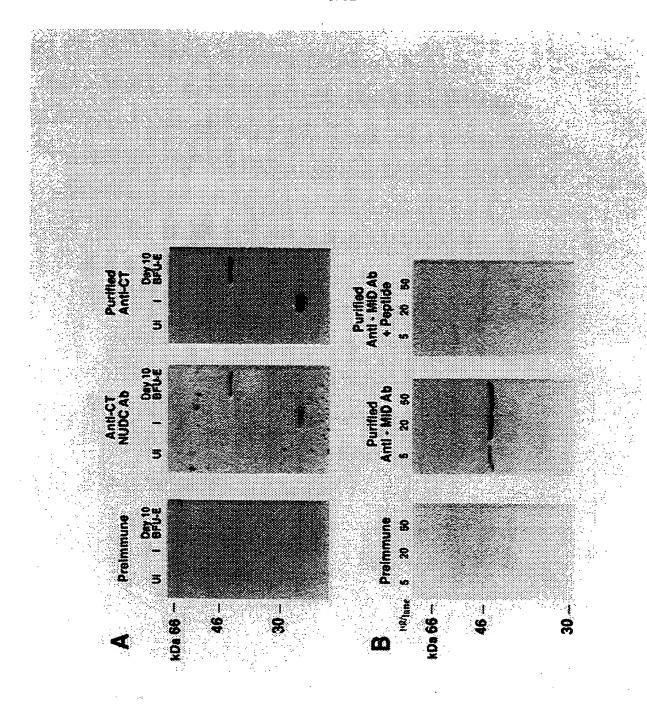
41 RAT LAKEAKAETPGPQIKELTDEEAERLQLEIDQKKDAENHEVQLKNGSLDSPGKQDAEEEEDEEKDKGKLKPNLGNGADLPNYRWTGTLS 180 131 ELDLAVPFCVNFRLKGKDM/VDIQRRHLRVGLKGQ--PAIIDGELYNEVKVEESSWLIED-----GKVVTVHLEKINKMEWWSRLVSS 260 ELDLAVPFRVSFRLKGKD WVDIQRRHLRVGLKGQ--APVIDGELYNEVKVEESSWLIED-----GKVVTVHLEKINKÆWWRLVTS 261 Lakeakset sgpoikeltdeeaerlole idokkdaenheaolkngsldspgkodtede-eedekdkgkikpnigngadlpnyrwtotls DVDVFTTI PVSANLKGRDLDVVLKKDS I KVKVKGENGEV FIDGQFPHP I KPSESSWTLETTSKPPFTGKEVS I HLDKVNQMEMWAHVVTT MSEQEPSSADLAAREAEEKQRK----AAEEAEQATLP-YKWTQTIR HL K N MEWW 202 HUM DPEINTKKINPENSKLSDLDSETRSMVERMMYDQRQKSMGLPTSDEQKKQEILKKFMDQHPEMDFSKAKFN 331 GK V **DPEINTKKINPENSKLSDLDSETRSMVEKMYDQRQKSMGLPTSDEQKKQEILKKEMDQHPEMDFSKAKFN** ASP APKIDVSKITPENSSLSDLDGETRAMVEKMANTDQRQKEMGAPTSDEQRKMDILKKFQKEHPEMDFSNAKIG HPEMDES AK ESSW KI PENS LSDLD ETR MVEKMMYDOROK MG PTSDEQ K ILKKF RAT RAT ASP CON CON

CON ASP



nudC cDNA pAL5 pAL5 + pAL5 + Hum. nudC Asp. nudC PAL5+

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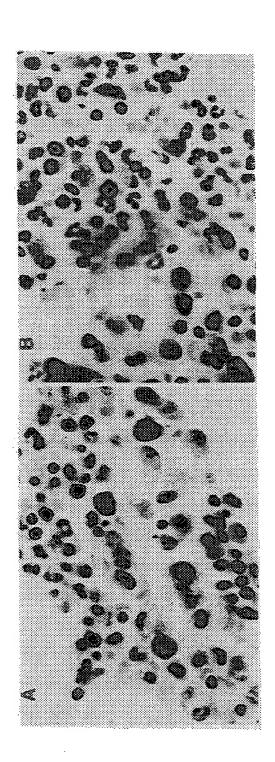


Fig. 4

PCT/US99/04996

5/12 .

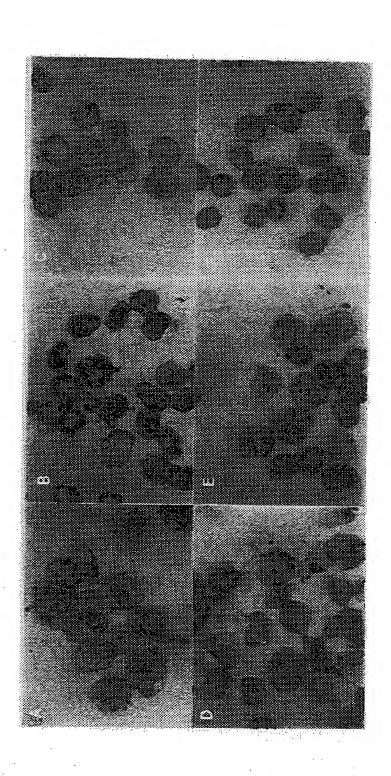


Fig. 5

PCT/US99/04996

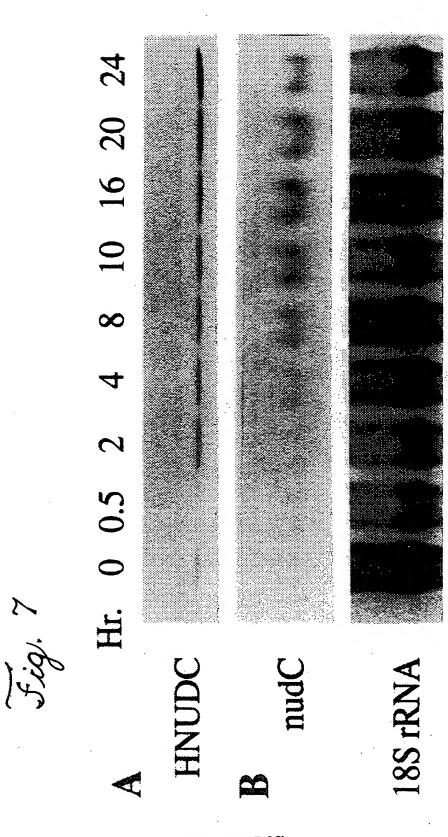
6/12

Fig. 6

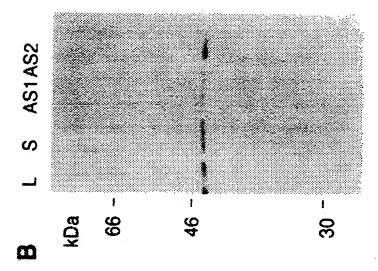
day 7 10 14 A

HNUDC -IκBα -

B
HnudC -



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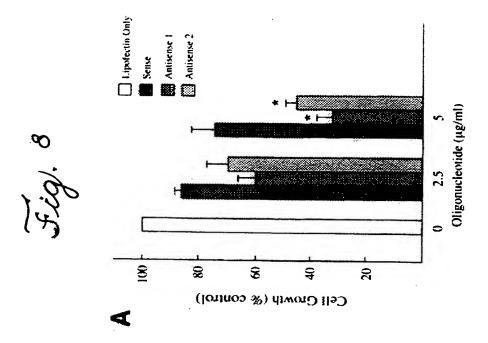




Fig.

HnudC Ribozyme: Con 2 3 4 6 2/3 2/4 2/6 3/4 3/6 4/6

Target RNA -

Fig. 10

FIG.11

1	CTAGAGTGCA GAGCTCCGGG ACGTGGATCG GAGCCGGCGC
	GATGGGCGGA GAGCAGGAGG
61	AGGAGCGGTT CGACGGCATG TTGCTGGCCA TGGCTCAGCA
	GCACGAGGGC GGCGTGCAGG
121	AGCTTGTGAA CACCTTCTTC AGCTTCCTTC GACGCAAAAC
	AGACTTTTC ATTGGAGGAG
181	AAGAAGGGAT GGCAGAGAAG CTTATCACAC AGACTTTCAG
	CCACCACAAT CAGCTGGCAC
241	AGAAGACCCG GCGGGAGAAG AGAGCCCGGC AGGAGGCCGA
	GCGGCGGAG AAGGCGGAGC
301	GGGCGGCCAG ACTGGCCAAG GAAGCCAAGT CAGAGACCTC
	AGGGCCCCAG ATCAAGGAGC
361	TAACTGATGA AGAGGCAGAG AGGCTGCAGC TAGAGATTGA
	CCAGAAAAG GATGCAGAGA
421	ATCATGAGGC CCAGCTCAAG AACGGCAGCC TTGACTCCCC
	AGGGAAGCAG GATACTGAGG
481	AAGATGAGGA GGAAGATGAG AAGGACAAAG GAAAACTGAA
	GCCCAACCTA GGCAACGGGG
541	CAGACCTGCC CAATTACCGC TGGACCCAGA CCCTGTCGGA
	GCTGGACCTG GCGGTCCCTT
601	TCTGTGTGAA CTTCCGGCTG AAAGGGAAGG ACATGGTGGT
	GGACATCCAG CGGCGCACC
661	TCCGGGTGGG GCTCAAGGGG CAGCCAGCGA TCATTGATGG
	GGAGCTCTAC AATGAAGTGA
721	AGGTGGAGGA GAGCTCGTGG CTCATTGAGG ACGGCAAGGT
	GGTGACTGTG CATCTGGAGA
781	AGATCAATAA GATGGAGTGG TGGAGCCGCT TGGTGTCCAG
	TGACCCTGAG ATCAACACCA







341	AGAAGATTAA CCCTGAGAAT TCCAAGCTGT CAGACCTGGA
	CAGTGAGACT CGCAGCATGG
901	TGGAAAAGAT GATGTATGAC CAGCGACAGA AGTCCATGGG
	GCTGCCAACT TCAGACGAAC
961	AGAAGAAACA GGAGATTCTG AAGAAGTTCA TGGATCAACA
	TCCGGAGATG GATTTTCCA
1021	AGGCTAAATT CAACTAGCCC CTGTTTTTTC CTCCCTGAAC
	TCTTGGGGCT GAGCTGCAAC
1081	CACCCAACTT TCTTTCCCAC TCTTCTCTGG GACTTGTGGG
	CCTCAGGGCT TGGGGCAGGC
1141	ATGGGACTGG CCCAGGCACA CAGGTCCCGG GGCATCAGGA
	GAAAGGCTGG GTCTTGGGAC
1201	CTTGTCCTCC CCAGTTGGCC TACTGTTACA CATTAAAACG
	ATTTGCCCAG CTCAAAAAA
1261	A A A A A A A A A A A A A A A A A A A

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